



SEQUENCE LISTING

<10> Glimcher, Laurie H.
Szabo, Susanne, J.

<120> T-BET COMPOSITIONS AND METHODS OF USE THEREOF

<130> HUI-040PC

<140> US 10/008/264

<141> 2001-12-03

<150> PCT/US00/17545

<151> 2000-06-01

<150> US 60/137085

<151> 1999-06-02

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<170> PatentIn Ver. 2.0

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<211> 1608

<212> DNA

<213> Homo sapiens

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<221> CDS

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Pro Met Pro Gly Ser Asp Glu Gly Arg Ala Pro Gly Ala Asp Pro Gln
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cac cgc tac ttc tac ccg gag ccg ggc gcg cag gac gcg gac gag cgt 144
His Arg Tyr Phe Tyr Pro Glu Pro Gly Ala Gln Asp Ala Asp Glu Arg
35 40 45

cgc ggg ggc ggc agc ctg ggg tct ccc tac ccg ggg ggc gcc ttg gtg 192
Arg Gly Gly Gly Ser Leu Gly Ser Pro Tyr Pro Gly Gly Ala Leu Val
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ccc gcc ccg ccg agc cgc ttc ctt gga gcc tac gcc tac ccg ccg cga 240
Pro Ala Pro Pro Ser Arg Phe Leu Gly Ala Tyr Ala Tyr Pro Pro Arg
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ccc cag gcg gcc ggc ttc ccc ggc gcg ggc gag tcc ttc ccg ccg ccc 288
Pro Gln Ala Ala Gly Phe Pro Gly Ala Gly Glu Ser Phe Pro Pro Pro
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gcg gac gcc gag ggc tac cag ccg ggc gag ggc tac gcc gcc ccg gac 336
Ala Asp Ala Glu Gly Tyr Gln Pro Gly Glu Gly Tyr Ala Ala Pro Asp
100 105 110

ccg cgc gcc ggg ctc tac ccg ggg ccg cgt gag gac tac gcg cta ccc 384

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gcg gga ctg gag gtg tcg ggg aaa ctg agg gtc gcg ctc aac aac cac	432
Ala Gly Leu Glu Val Ser Gly Lys Leu Arg Val Ala Leu Asn Asn His	
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ctg ttg tgg tcc aag ttt aat cag cac cag aca gag atg atc atc acc	480
Leu Leu Trp Ser Lys Phe Asn Gln His Gln Thr Glu Met Ile Ile Thr	
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aag cag gga cgg cgg atg ttc cca ttc ctg tca ttt act gtg gcc ggg	528
Lys Gln Gly Arg Arg Met Phe Pro Phe Leu Ser Phe Thr Val Ala Gly	
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ctg gag ccc acc agc cac tac agg atg ttt gtg gac gtg gtc ttg gtg	576
Leu Glu Pro Thr Ser His Tyr Arg Met Phe Val Asp Val Val Leu Val	
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gac cag cac cac tgg cgg tac cag agc ggc aag tgg gtg cag tgt gga	624
Asp Gln His His Trp Arg Tyr Gln Ser Gly Lys Trp Val Gln Cys Gly	
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aag gcc gag ggc agc atg cca gga aac cgc ctg tac gtc cac ccg gac	672
Lys Ala Glu Gly Ser Met Pro Gly Asn Arg Leu Tyr Val His Pro Asp	
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Val Glu Val Asn Asp Gly Glu Pro Glu Ala Ala Cys Asn Ala Ser Asn	
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acg cat atc ttt act ttc caa gaa acc cag ttc att gcc gtg act gcc	912
Thr His Ile Phe Thr Phe Gln Glu Thr Gln Phe Ile Ala Val Thr Ala	
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Tyr Gln Asn Ala Glu Ile Thr Gln Leu Lys Ile Asp Asn Asn Pro Phe	
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Ala Lys Gly Phe Arg Glu Asn Phe Glu Ser Met Tyr Thr Ser Val Asp	
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acc agc atc ccc tcc ccg cct gga ccc aac tgt caa ttc ctt ggg gga	1056
Thr Ser Ile Pro Ser Pro Pro Gly Asn Cys Gln Phe Leu Gly Gly	
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Tyr Trp Leu Gly Ala Pro Arg Asp His Ser Tyr	Glu Ala Glu Phe Arg		
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Met Ser Tyr Tyr Arg Gly Gln Glu Val Leu Ala Pro	Gly Ala Gly Trp		
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Pro Val Ala Pro Gln Tyr Pro Pro Lys Met Gly Pro Ala	Ser Trp Phe		
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Arg Pro Met Arg Thr Leu Pro Met Glu Pro Gly Pro Gly	Gly Ser Glu		
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gcc ccc atc cgg ccg gaa tcc agt gat tca gga ctg ggc	gaa gga gac	1488	
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Ser Lys Arg Arg Arg Val Ser Pro Tyr Pro Ser Ser Gly	Asp Ser Ser		
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<212> PRT

<213> Homo sapiens

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Arg Gly Gly Gly Ser Leu Gly Ser Pro Tyr Pro Gly Gly Ala Leu Val
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Pro Ala Pro Pro Ser Arg Phe Leu Gly Ala Tyr Ala Tyr Pro Pro Arg
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Pro Gln Ala Ala Gly Phe Pro Gly Ala Gly Glu Ser Phe Pro Pro Pro
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Pro Arg Ala Gly Leu Tyr Pro Gly Pro Arg Glu Asp Tyr Ala Leu Pro
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Ala Gly Leu Glu Val Ser Gly Lys Leu Arg Val Ala Leu Asn Asn His
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Leu Leu Trp Ser Lys Phe Asn Gln His Gln Thr Glu Met Ile Ile Thr
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Lys Gln Gly Arg Arg Met Phe Pro Phe Leu Ser Phe Thr Val Ala Gly
165 170 175

Leu Glu Pro Thr Ser His Tyr Arg Met Phe Val Asp Val Val Leu Val
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Asp Gln His His Trp Arg Tyr Gln Ser Gly Lys Trp Val Gln Cys Gly
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Lys Ala Glu Gly Ser Met Pro Gly Asn Arg Leu Tyr Val His Pro Asp
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Ser Pro Asn Thr Gly Ala His Trp Met Arg Gln Glu Val Ser Phe Gly
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Lys Leu Lys Leu Thr Asn Asn Lys Gly Ala Ser Asn Asn Val Thr Gln
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Met Ile Val Leu Gln Ser Leu His Lys Tyr Gln Pro Arg Leu His Ile
260 265 270

Val Glu Val Asn Asp Gly Glu Pro Glu Ala Ala Cys Asn Ala Ser Asn
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Thr His Ile Phe Thr Phe Gln Glu Thr Gln Phe Ile Ala Val Thr Ala
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Tyr Gln Asn Ala Glu Ile Thr Gln Leu Lys Ile Asp Asn Asn Pro Phe
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Asp His Tyr Ser Pro Leu Leu Pro Asn Gln Tyr Pro Val Pro Ser Arg
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 385 390 395 400
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 Met Ser Tyr Tyr Arg Gly Gln Glu Val Leu Ala Pro Gly Ala Gly Trp
 420 425 430
 Pro Val Ala Pro Gln Tyr Pro Pro Lys Met Gly Pro Ala Ser Trp Phe
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 Arg Pro Met Arg Thr Leu Pro Met Glu Pro Gly Pro Gly Gly Ser Glu
 450 455 460
 Gly Arg Gly Pro Glu Asp Gln Gly Pro Pro Leu Val Trp Thr Glu Ile
 465 470 475 480
 Ala Pro Ile Arg Pro Glu Ser Ser Asp Ser Gly Leu Gly Glu Gly Asp
 485 490 495
 Ser Lys Arg Arg Arg Val Ser Pro Tyr Pro Ser Ser Gly Asp Ser Ser
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 Pro Met Pro Ser Asp Glu Gly Arg Gly Pro Gly Ala Asp Gln Gln His
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 Arg Phe Phe Tyr Pro Glu Pro Gly Ala Gln Asp Pro Thr Asp Arg Arg
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 Ala Gly Ser Ser Leu Gly Thr Pro Tyr Ser Gly Gly Ala Leu Val Pro
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 gcc gcg ccg ggt cgc ttc ctt gga tcc ttc gcc tac ccg ccc cgg gct 240
 Ala Ala Pro Gly Arg Phe Leu Gly Ser Phe Ala Tyr Pro Pro Arg Ala

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ggt gcg gag ggc tac ccg ccc gtg gat ggc tac cct gcc cct gac ccg Gly Ala Glu Gly Tyr Pro Pro Val Asp Gly Tyr Pro Ala Pro Asp Pro	100	105	336
cgc gcg ggg ctc tac cca ggg ccg cgc gag gac tac gca ttg ccc gcg Arg Ala Gly Leu Tyr Pro Gly Pro Arg Glu Asp Tyr Ala Leu Pro Ala	115	120	384
ggg ttg gag gtg tct ggg aag ctg aga gtc gcg ctc agc aac cac ctg Gly Leu Glu Val Ser Gly Lys Leu Arg Val Ala Leu Ser Asn His Leu	130	135	432
ttg tgg tcc aag ttc aac cag cac cag aca gag atg atc atc act aag Leu Trp Ser Lys Phe Asn Gln His Gln Thr Glu Met Ile Ile Thr Lys	145	150	480
caa gga cgg cga atg ttc cca ttc ctg tcc ttc acc gtg gcc ggg ctg Gln Gly Arg Arg Met Phe Pro Phe Leu Ser Phe Thr Val Ala Gly Leu	165	170	528
gag ccc aca agc cat tac agg atg ttt gtg gat gtg gtc ttg gtg gac Glu Pro Thr Ser His Tyr Arg Met Phe Val Asp Val Val Leu Val Asp	180	185	576
cag cac cac tgg cgg tac cag agc ggc aag tgg gtg cag tgt gga aag Gln His His Trp Arg Tyr Gln Ser Gly Lys Trp Val Gln Cys Gly Lys	195	200	624
gca gaa ggc agc atg cca ggg aac cgc tta tat gtc cac cca gac tcc Ala Glu Gly Ser Met Pro Gly Asn Arg Leu Tyr Val His Pro Asp Ser	210	215	672
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cag aac gca gag atc act cag ctg aaa atc gac aac aac ccc ttt gcc Gln Asn Ala Glu Ile Thr Gln Leu Lys Ile Asp Asn Asn Pro Phe Ala	305	310	960
		315	320

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 Lys Gly Phe Arg Glu Asn Phe Glu Ser Met Tyr Ala Ser Val Asp Thr
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agt gtc ccc tcg cca cct gga ccc aac tgt caa ctg ctt ggg gga gac 1056
 Ser Val Pro Ser Pro Pro Gly Pro Asn Cys Gln Leu Leu Gly Gly Asp
 340 345 350

ccc ttc tca cct ctt cta tcc aac cag tat cct gtt ccc agc cgt ttc 1104
 Pro Phe Ser Pro Leu Leu Ser Asn Gln Tyr Pro Val Pro Ser Arg Phe
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 Tyr Pro Asp Leu Pro Gly Gln Pro Lys Asp Met Ile Ser Gln Pro Tyr
 370 375 380

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 Trp Leu Gly Thr Pro Arg Glu His Ser Tyr Glu Ala Glu Phe Arg Ala
 385 390 395 400

gtg agc atg aag ccc aca ctc cta ccc tct gcc ccg ggg ccc act gtg 1248
 Val Ser Met Lys Pro Thr Leu Leu Pro Ser Ala Pro Gly Pro Thr Val
 405 410 415

ccc tac tac cgg ggc caa gac gtc ctg gcg cct gga gct ggt tgg ccc 1296
 Pro Tyr Tyr Arg Gly Gln Asp Val Leu Ala Pro Gly Ala Gly Trp Pro
 420 425 430

gtg gcc cct caa tac ccg ccc aag atg agc cca gct ggc tgg ttc cgg 1344
 Val Ala Pro Gln Tyr Pro Pro Lys Met Ser Pro Ala Gly Trp Phe Arg
 435 440 445

ccc atg cga act ctg ccc atg gac ccg ggc ctg gga tcc tca gag gaa 1392
 Pro Met Arg Thr Leu Pro Met Asp Pro Gly Leu Gly Ser Ser Glu Glu
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cag ggc tcc tcc ccc tcg ctg tgg cct gag gtc acc tcc ctc cag ccg 1440
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 Glu Pro Ser Asp Ser Gly Leu Gly Glu Gly Asp Thr Lys Arg Arg Arg
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ata tcc ccc tat cct tcc agt ggc gac agc tcc tct ccc gct ggg gcc 1536
 Ile Ser Pro Tyr Pro Ser Ser Gly Asp Ser Ser Ser Pro Ala Gly Ala
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cct tct cct ttt gat aag gaa acc gaa ggc cag ttt tat aat tat ttt 1584
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 Ala Gly Ser Ser Leu Gly Thr Pro Tyr Ser Gly Gly Ala Leu Val Pro
 50 55 60
 Ala Ala Pro Gly Arg Phe Leu Gly Ser Phe Ala Tyr Pro Pro Arg Ala
 65 70 75 80
 Gln Val Ala Gly Phe Pro Gly Pro Gly Glu Phe Phe Pro Pro Pro Ala
 85 90 95
 Gly Ala Glu Gly Tyr Pro Pro Val Asp Gly Tyr Pro Ala Pro Asp Pro
 100 105 110
 Arg Ala Gly Leu Tyr Pro Gly Pro Arg Glu Asp Tyr Ala Leu Pro Ala
 115 120 125
 Gly Leu Glu Val Ser Gly Lys Leu Arg Val Ala Leu Ser Asn His Leu
 130 135 140
 Leu Trp Ser Lys Phe Asn Gln His Gln Thr Glu Met Ile Ile Thr Lys
 145 150 155 160
 Gln Gly Arg Arg Met Phe Pro Phe Leu Ser Phe Thr Val Ala Gly Leu
 165 170 175
 Glu Pro Thr Ser His Tyr Arg Met Phe Val Asp Val Val Leu Val Asp
 180 185 190
 Gln His His Trp Arg Tyr Gln Ser Gly Lys Trp Val Gln Cys Gly Lys
 195 200 205
 Ala Glu Gly Ser Met Pro Gly Asn Arg Leu Tyr Val His Pro Asp Ser
 210 215 220
 Pro Asn Thr Gly Ala His Trp Met Arg Gln Glu Val Ser Phe Gly Lys
 225 230 235 240
 Leu Lys Leu Thr Asn Asn Lys Gly Ala Ser Asn Asn Val Thr Gln Met
 245 250 255
 Ile Val Leu Gln Ser Leu His Lys Tyr Gln Pro Arg Leu His Ile Val
 260 265 270
 Glu Val Asn Asp Gly Glu Pro Glu Ala Ala Cys Ser Ala Ser Asn Thr
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Lys Gly Phe Arg Glu Asn Phe Glu Ser Met Tyr Ala Ser Val Asp Thr
325 330 335

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Tyr Pro Asp Leu Pro Gly Gln Pro Lys Asp Met Ile Ser Gln Pro Tyr
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Trp Leu Gly Thr Pro Arg Glu His Ser Tyr Glu Ala Glu Phe Arg Ala
385 390 395 400

Val Ser Met Lys Pro Thr Leu Leu Pro Ser Ala Pro Gly Pro Thr Val
405 410 415

Pro Tyr Tyr Arg Gly Gln Asp Val Leu Ala Pro Gly Ala Gly Trp Pro
420 425 430

Val Ala Pro Gln Tyr Pro Pro Lys Met Ser Pro Ala Gly Trp Phe Arg
435 440 445

Pro Met Arg Thr Leu Pro Met Asp Pro Gly Leu Gly Ser Ser Glu Glu
450 455 460

Gln Gly Ser Ser Pro Ser Leu Trp Pro Glu Val Thr Ser Leu Gln Pro
465 470 475 480

Glu Pro Ser Asp Ser Gly Leu Gly Glu Gly Asp Thr Lys Arg Arg Arg
485 490 495

Ile Ser Pro Tyr Pro Ser Ser Gly Asp Ser Ser Ser Pro Ala Gly Ala
500 505 510

Pro Ser Pro Phe Asp Lys Glu Thr Glu Gly Gln Phe Tyr Asn Tyr Phe
515 520 525

Pro Asn
530